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FIG. 1

DNA (TRANSCRIPTION) - MRNA (TRANSLATION) - PROTEIN

3000000 I TRANSLATION HALTED #RNA HYBRID OLIGONUCL. DIRECTION OF TRANSLATION POLYRIBOSOME DWA FOR T PROTEIN [

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FIG. 3 OLIGONUCLEOTIDE SPECIFIC FOR SV-40 T PROTEIN VIRAL DNA(CODING) 3'... GAAA CGTTTCTACCTATTTC...5' VIRAL T PROTEIN MRNA 5'... CUUUG CAAAGAUGGAUAAAG...3' T PROTEIN SPECIFIC OLIGONUCLEOTIDE 3'... GAAA CGTTTCTACCTATTTC...5'

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LINKER T PROTEIN OLIGONUCLEOTIDE CTAAGCTTAG GATTCGAATC 7-4 DNA LIGASE TINKER CONSTRUCTION OF HIGH YIELD PLASMID CONTAINIING THE T PROTEIN OLIGONUCLEOTIDE ACTAAGCTTAGCTTTATCCATCTTTGCAAAGCTAAGCTTAGT PCTTTATCCATCTTTGCAAAG GAAATAGGTAGAAACGTTTCP 7-411GASE PT PROTEIN OLIGONUCLEOTIDE 1 (Tcs, Apr) (ONE STRAND SHOWN) SV40 T PROTEIN OLIGONUCLEOTIDE (DOUBLE STRANDED) SI NUCLEASE (REMOVE COHESIVE ENDS) HIND III NUCLEASE (CLEAVES IN TO GENE) COVALENTLY CLOSED CIRCLE) P BR 312 (CHOPPED)

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FIG.5

PRODUCTION OF SV-40 T PROTEIN OLIGONUCLEOTIDE FROM PT OLIGONUCLEOTIDE USING RESTRICTION NUCLEASE CLEAVAGE TO RELEASE FRAGMENTS.

ACTAAGCTTAGCTTTATCCATCTTTGCAAAGCTAAGCTTAGT PT PROTEIN OLIGONUCLEOTIDE 1 (ONE STRAND CODE SHOWN) RESTRICTION NUCLEASE HINDIII (CLEAVES AT AAGCTT) ACTA | AGCTTAGCTTTATCCATACTTTGCAAAGCTA | AGCTTAGT TGATTCGA | ATCGAAATAGGTATGAMCGTTTCGATTCGA | ATCA AGCTTAGCTTTATCCATCTTTGCAAAGCTA AT CGAAATAGGTAGAAACGTTTCGATTCGA PBR322 FRAGMENTS (REMOVED BY GEL FILTRATION) LINKEREEIN OLGONUCTLIEE-OIDNKLRE A U I RTEREST I ON N ULCE A SE VORE SAG)T A +DIAUSSTREOMOVE ENDF AMBTS CTT TA C CA T GT T AGA GAA AT GAG AG A AGCCT TT TPRTO EOGNN OEW & D PRO CE BOS MKA BIGN BL BID OLG OBE OT IDE I A PTHROTSERS O CT TA T CACTICE AGA A

T-PROETI ON LG BUCLEO IDE (POLA HOSTPHER) RIEST Docket No.: ISIS-4502

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FIG.6

DNA SEGMENT

ACETY DATE BEAVSA OPTEE TO T

RTERB DN A PO L YSPAHADO

DNA PODSPHOTEIES

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FIG.7

OLIGONUCLEOTIDE SPECIFIC FOR FOLLICLE STIMULATING HORMONE

PROTEIN SEQUENCE:

N TERMINUS THR TRP CYS ALA GLY TYR CYS TYR THR ... C TERMINUS

MRNA PREDICTED:
5'END ACX UGG UGC GCX GGX VAC UGU UAC ACX 3'END

OLIGONUCLEOTIDE

OLIGONUCLEOTIDE

FAMILY:

3' END TCG ACC ACC CGG CCG ATG ACG ATG TG- 5' END

(f)

(f)